Revising the chromosome-specific probes of white hawk (Leucopternis albicollis)

Ivanete Furo ¹, Rafael Kretschmer ², Jorge Pereira ³, Darren Griffin ⁴, Rebecca O'Connor ⁴, Patricia O'Brien ⁵, Malcolm Ferguson-Smith ⁵, Edivaldo de Oliveira ^{6,7}

¹Laboratório de Reprodução Animal, LABRAC, Universidade Federal Rural da Amazônia, UFRA, Parauapebas, PA, Brazil;

² Programa de Pós-graduação em Genética e Biologia Molecular, PPGBM, Universidade Federal do Rio Grande do Sul, Porto Alegre, RS, Brazil;

³ Animal and Veterinary Research Centre (CECAV), University of Trás-os-Montes and Alto Douro (UTAD), Vila Real, Portugal.

⁴ School of Biosciences, University of Kent, Canterbury CT2 7NJ, UK;

⁵ Cambridge Resource Centre for Comparative Genomics, University of Cambridge Department of Veterinary Medicine, Cambridge, United Kingdom;

⁶ Laboratório de Cultura de Tecidos e Citogenética, SAMAM, Instituto Evandro Chagas, Ananindeua, PA, Brazil.

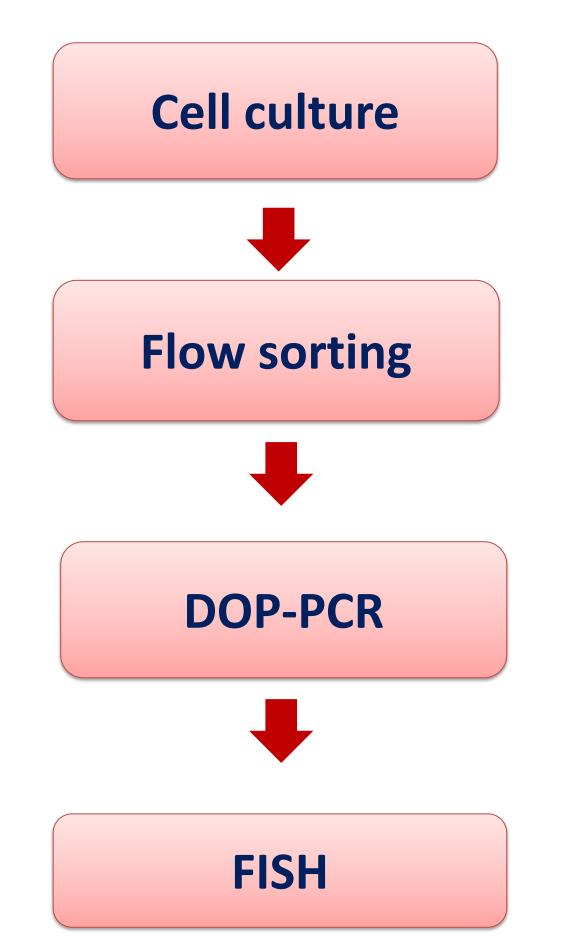
⁷ Instituto de Ciências Exatas e Naturais, Universidade Federal do Pará, Belém-PA-Brazil;



Introduction

Leucopternis albicollis is a diurnal bird of prey with extensive karyotype reorganization. Chromosome-specific probes from this species have been used successfully to detect intrachromosomal rearrangements in different species of bird since 2010. However, some gaps were detected using probes obtained from the first set. Here, we have obatined a new set of whole chromosome probes in order to improve the previous one; also we have performed experiments using bacterial artificial chromosome (BAC) from chicken microchromosomes.

Methods and Results



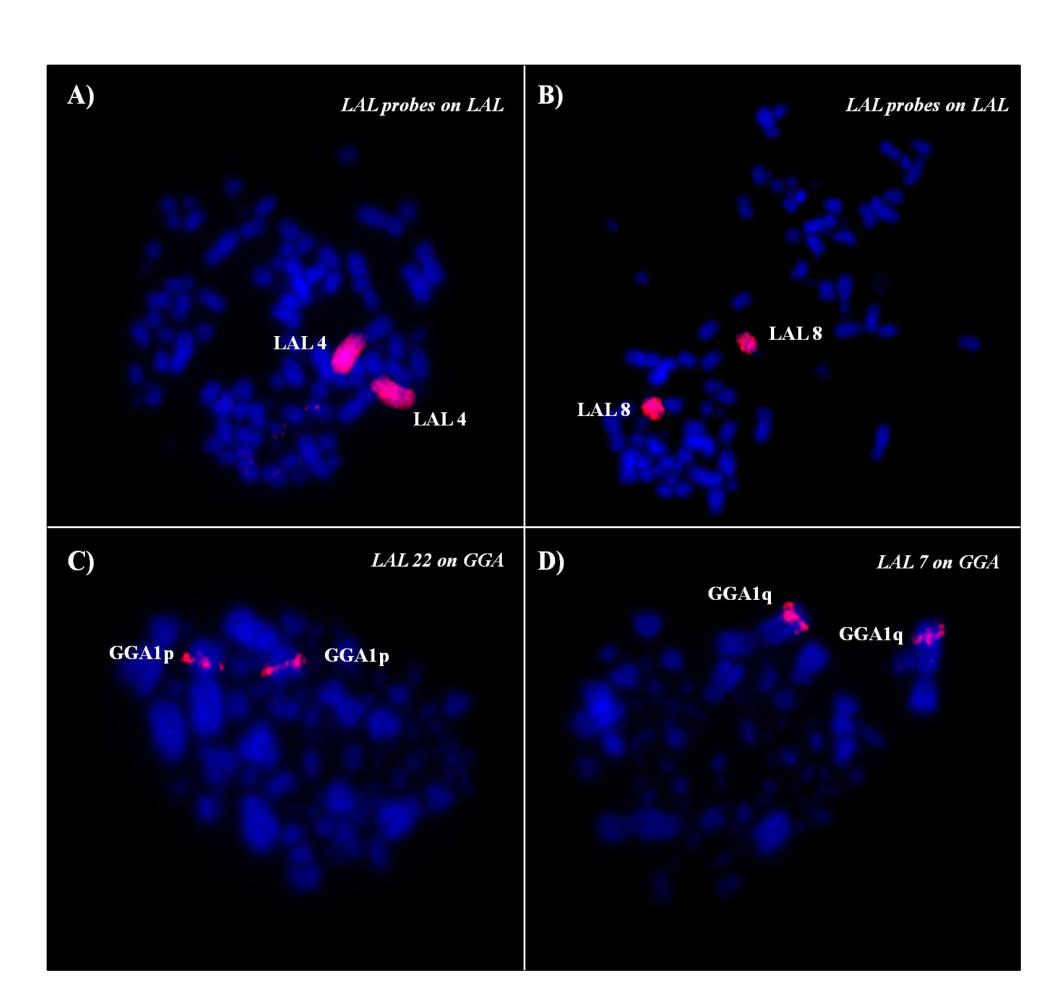


Figure.1 Representative Same-species FISH experiments using *L. albicollis* (LAL) probes: LAL 4 (A), LAL8 (B) and Cross-species LAL probes on *G.gallus* metaphases (C and D)

Conclusion

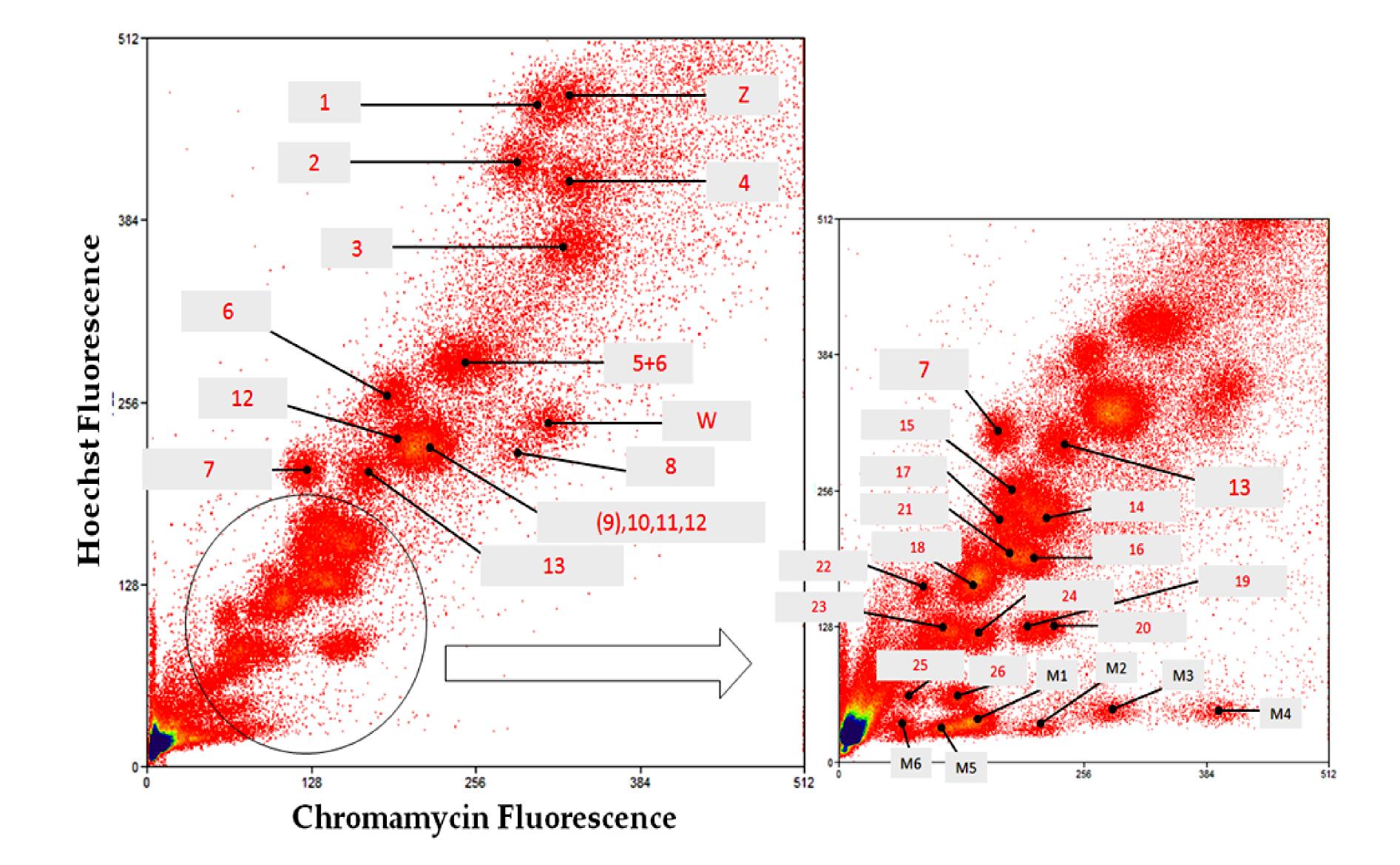


Figure.2 New flow karyotype of *L. albicollis*. Legend: M= Microchromosomes.

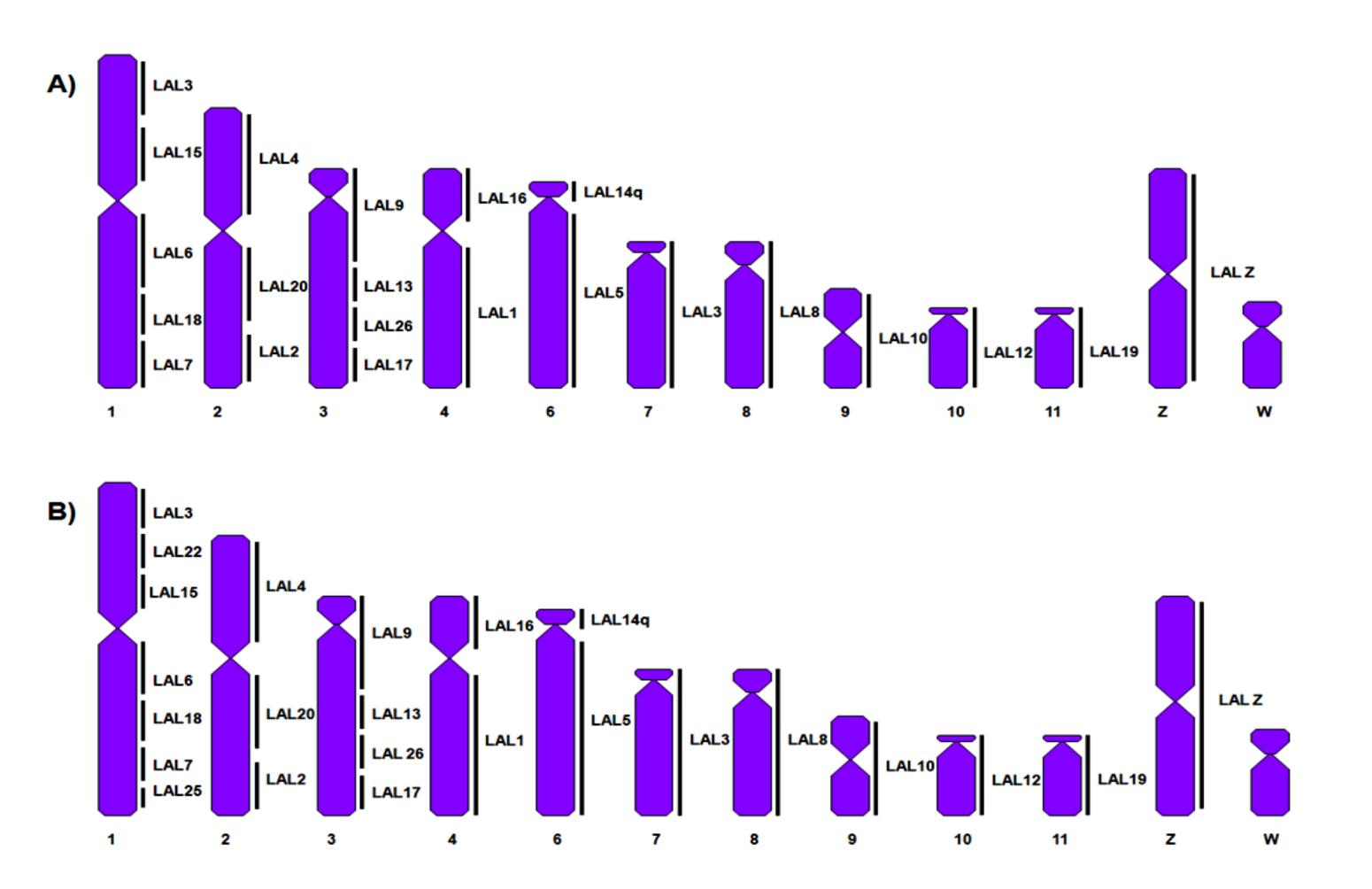


Figure.3 Homology map between chicken macrochromosomes and white hawk paints. (A) The homology described by de Oliveira et al. [4] (B) The new nomenclature has been proposed for the new set of probes from white hawk.

Our results show that microchromosomes 17-28 were involved in fusion events with macrochromosomes. In addition, a new nomenclature has been proposed for the new set of probes and some previous inaccuracies corrected. As an example, ancestral chromosome GGA1 is now shown to have homology to seven white hawk chromosomes rather than five, as homologies to two small chromosomes were missed. In conclusion, the new complete set of chromosome probes will improve the value of this tool for avian comparative cytogenetics.